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Protein structure and disorder analysis of the first 120 amino acids of the protein. The analysis is divided into three segments: 1-60, 70-120, and 130-150. Each segment shows the sequence, secondary structure (alpha-helices in green, beta-strands in blue), solvent accessibility (SS), confidence scores, and disorder predictions. The first 60 amino acids show a high degree of disorder, while the subsequent segments show more structured regions.

Confidence Key

High(9)  Low (0)

? Disordered

 Alpha helix

 Beta strand